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Genomic sequencing of human chromosome 19 and comparative analysis of human and rodent DNA repair gene regions.

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Approximately 70% of chromosome 19 is spanned by cosmids for which an EcoR1 map has been derived. We are now scaling our sequencing efforts to take advantage of these pre-mapped clones. To date, we have completed ~500 kbp of genomic sequence which has been targeted primarily to cosmids containing the human DNA repair genes *XRCC1* and *ERCC2* on chromosome 19, *ERCC4* on chromosome 16, and *XRCC3* on chromosome 14, as well as selected rodent homologs. We have sequenced 76 kbp containing the human and mouse *XRCC1* genes, which span 26 kbp in the mouse and 31.9 kbp in the human. In addition to the coding regions, 9 conserved elements were identified with sequence identities ranging from 65% to 78%. We have completed 52 kbp of human sequence encompassing the *ERCC2* gene as well as 54 kb spanning the syntenic regions in the mouse and hamster. A defect in *ERCC2* leads to the cancer-prone human disorder xeroderma pigmentosum (XP-D). The human *ERCC2* gene is comprised of 23 exons and is 98% identical to the rodent homologs at the protein level. We identified two genes flanking *ERCC2*, one may be a new member of the kinesin gene family, and the other has no known function. Like *ERCC2*, the *ERCC4* gene product is involved in the nucleotide excision repair pathway, which recognizes and removes DNA damage. A total of 35 kbp has been completed for this gene region which has been instrumental in identifying and assembling the coding regions from numerous partial length cDNAs. The full-length gene spans ~29 kbp and is >50% AT-rich. The *ERCC4* gene product exhibits significant homology to the *S. cerevisiae* rad1 and *S. pombe* rad16 genes, which encode single strand endonucleases. Finally, we have completed sequencing a 2.7 kbp candidate cDNA for the recently cloned human *XRCC3* gene and are in the process of sequencing the cosmid containing this gene, which appears to play a crucial role in chromosomal stability. This work was performed by Lawrence Livermore National Laboratory under the auspices of the U.S. Department of Energy, Contract No. W-7405-Eng-48.